

chain nodes :

7 8 9 10 11 12 16 22 23 24 25 26 27 28 29 30 31 32 33 35

ring nodes :

1 2 3 4 5 6 17 18 19 20 21

chain bonds :

1-21 2-7 4-8 5-32 6-31 8-9 8-16 10-12 10-11 17-22 17-29 18-23 18-30 19-24

19-25 21-35 24-27 25-26 26-33 27-28

ring bonds :

1-2 1-6 2-3 3-4 4-5 5-6 17-18 17-21 18-19 19-20 20-21

exact/norm bonds :

1-2 1-6 1-21 2-3 2-7 3-4 4-5 4-8 5-6 8-16 10-12 10-11 17-18 17-21 17-22

18-19 18-23 19-20 19-24 20-21 24-27 26-33 27-28

exact bonds :

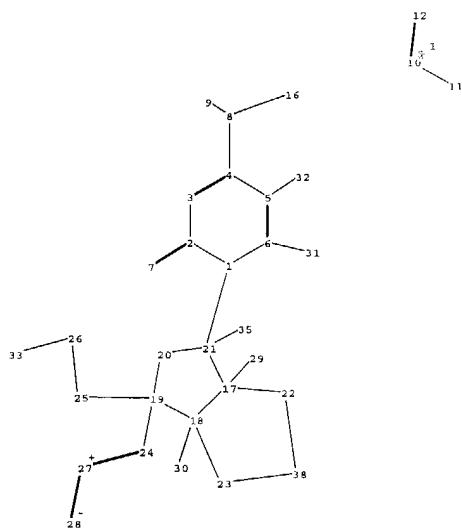
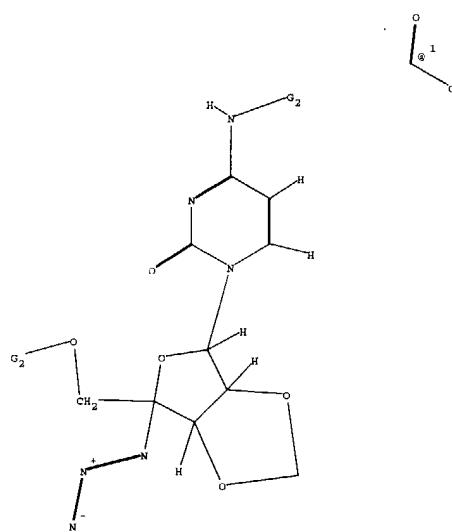
5-32 6-31 8-9 17-29 18-30 19-25 21-35 25-26

G1:O,S,N,C

G2:H,[*1]

Match level :

 1:Atom 2:Atom 3:Atom 4:Atom 5:Atom 6:Atom 7:CLASS 8:CLASS 9:CLASS 10:CLASS
 11:CLASS 12:CLASS 16:CLASS 17:Atom 18:Atom 19:Atom 20:Atom 21:Atom 22:CLASS
 23:CLASS 24:CLASS 25:CLASS 26:CLASS 27:CLASS 28:CLASS 29:CLASS 30:CLASS 31:CLASS
 32:CLASS 33:CLASS 35:CLASS



chain nodes :

7 8 9 10 11 12 16 24 25 26 27 28 29 30 31 32 33 35

ring nodes :

1 2 3 4 5 6 17 18 19 20 21 22 23 38

chain bonds :

1-21 2-7 4-8 5-32 6-31 8-9 8-16 10-12 10-11 17-29 18-30 19-24 19-25 21-35
24-27 25-26 26-33 27-28

ring bonds :

1-2 1-6 2-3 3-4 4-5 5-6 17-18 17-21 17-22 18-19 18-23 19-20 19-21 22-38 23-38

exact/norm bonds :

1-2 1-6 1-21 2-3 2-7 3-4 4-5 4-8 5-6 8-16 10-12 10-11 17-18 17-21 17-22
18-19 18-23 19-20 19-24 20-21 22-38 23-38 24-27 26-33 27-28

exact bonds :

5-32 6-31 8-9 17-29 18-30 19-25 21-35 25-26

G1:O,S,N,C

G2:H,[*1]

Match level :

1:Atom 2:Atom 3:Atom 4:Atom 5:Atom 6:Atom 7:CLASS 8:CLASS 9:CLASS 10:CLASS
11:CLASS 12:CLASS 16:CLASS 17:Atom 18:Atom 19:Atom 20:Atom 21:Atom 22:CLASS
23:CLASS 24:CLASS 25:CLASS 26:CLASS 27:CLASS 28:CLASS 29:CLASS 30:CLASS 31:CLASS
32:CLASS 33:CLASS 35:CLASS 38:Atom

=> d his

(FILE 'HOME' ENTERED AT 18:00:44 ON 10 AUG 2004)

FILE 'REGISTRY' ENTERED AT 18:00:56 ON 10 AUG 2004

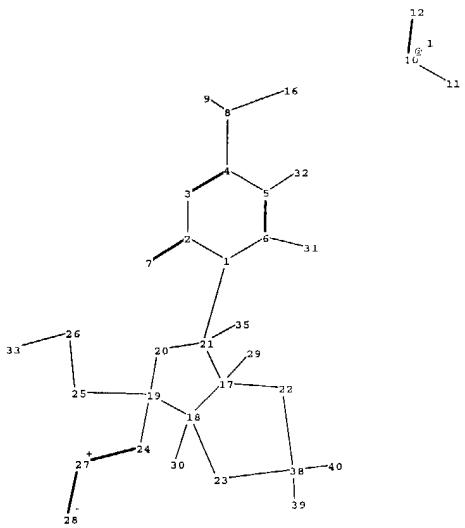
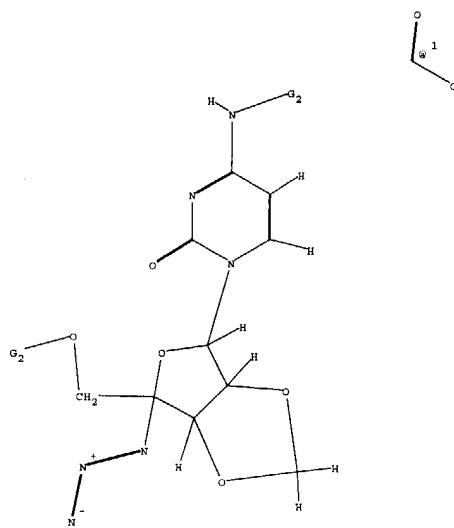
L1 STRUCTURE uploaded

L2 1 S L1 SSS SAM

L3 8 S L1 SSS FULL

FILE 'CAPLUS, MEDLINE' ENTERED AT 18:02:57 ON 10 AUG 2004

L4 1 S L3



chain nodes :

7 8 9 10 11 12 16 24 25 26 27 28 29 30 31 32 33 35 39 40

ring nodes :

1 2 3 4 5 6 17 18 19 20 21 22 23 38

chain bonds :

1-21 2-7 4-8 5-32 6-31 8-9 8-16 10-12 10-11 17-29 18-30 19-24 19-25 21-35
24-27 25-26 26-33 27-28 38-39 38-40

ring bonds :

1-2 1-6 2-3 3-4 4-5 5-6 17-18 17-21 17-22 18-19 18-23 19-20 20-21 22-38 23-38

exact/norm bonds :

1-2 1-6 1-21 2-3 2-7 3-4 4-5 4-8 5-6 8-16 10-12 10-11 17-18 17-21 17-22
18-19 18-23 19-20 19-24 20-21 22-38 23-38 24-27 26-33 27-28

exact bonds :

5-32 6-31 8-9 17-29 18-30 19-25 21-35 25-26 38-39 38-40

G1:O,S,N,C

G2:H,[*1]

Match level :

1:Atom 2:Atom 3:Atom 4:Atom 5:Atom 6:Atom 7:CLASS 8:CLASS 9:CLASS 10:CLASS
11:CLASS 12:CLASS 16:CLASS 17:Atom 18:Atom 19:Atom 20:Atom 21:Atom 22:CLASS
23:CLASS 24:CLASS 25:CLASS 26:CLASS 27:CLASS 28:CLASS 29:CLASS 30:CLASS 31:CLASS
32:CLASS 33:CLASS 35:CLASS 38:Atom 39:CLASS 40:CLASS

=> s 11 sss sam
SAMPLE SEARCH INITIATED 18:19:22 FILE 'REGISTRY'
SAMPLE SCREEN SEARCH COMPLETED - 15 TO ITERATE

100.0% PROCESSED 15 ITERATIONS 0 ANSWERS
SEARCH TIME: 00.00.01

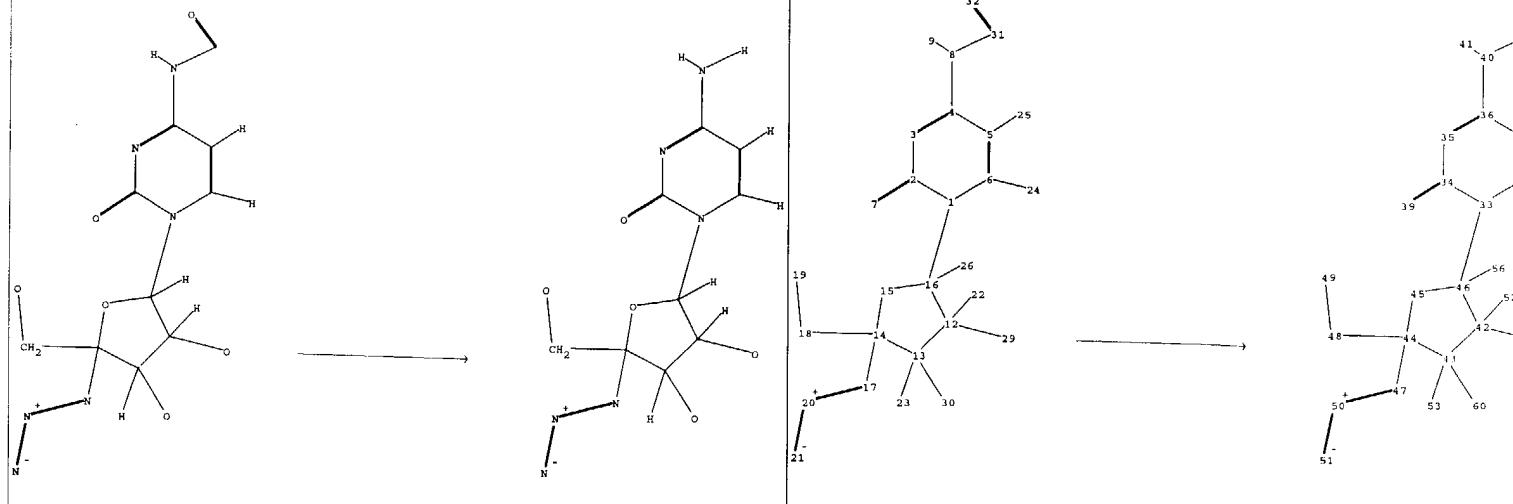
FULL FILE PROJECTIONS: ONLINE **COMPLETE**
BATCH **COMPLETE**
PROJECTED ITERATIONS: 68 TO 532
PROJECTED ANSWERS: 0 TO 0

L2 0 SEA SSS SAM L1

=> s 11 sss full
FULL SEARCH INITIATED 18:19:36 FILE 'REGISTRY'
FULL SCREEN SEARCH COMPLETED - 221 TO ITERATE

100.0% PROCESSED 221 ITERATIONS 0 ANSWERS
SEARCH TIME: 00.00.01

L3 0 SEA SSS FUL L1



chain nodes :

7	8	9	17	18	19	20	21	22	23	24	25	26	29	30	31	32	39	40	41	47	48	49	50
51	52	53	54	55	56	59	60	61															

ring nodes :

1	2	3	4	5	6	12	13	14	15	16	33	34	35	36	37	38	42	43	44	45	46	
---	---	---	---	---	---	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	----	--

chain bonds :

1-16	2-7	4-8	5-25	6-24	8-9	8-31	12-22	12-29	13-23	13-30	14-17	14-18	16-26								
17-20	18-19	20-21	31-32	33-46	34-39	36-40	37-55	38-54	40-41	40-61	42-52	42-59									
43-53	43-60	44-47	44-48	46-56	47-50	48-49	50-51														

ring bonds :

1-2	1-6	2-3	3-4	4-5	5-6	12-13	12-16	13-14	14-15	15-16	33-34	33-38	34-35	35-36							
36-37	37-38	42-43	42-46	43-44	44-45	45-46															

exact/norm bonds :

1-2	1-6	1-16	2-3	2-7	3-4	4-5	4-8	5-6	8-31	12-13	12-16	12-29	13-14	13-30							
14-15	14-17	15-16	17-20	20-21	31-32	33-34	33-38	33-46	34-35	34-39	35-36	36-37									
36-40	37-38	42-43	42-46	42-59	43-44	43-60	44-45	44-47	45-46	47-50	50-51										

exact bonds :

5-25	6-24	8-9	12-22	13-23	14-18	16-26	18-19	37-55	38-54	40-41	40-61	42-52									
43-53	44-48	46-56	48-49																		

G1:O,S,N,C

G2:H

Match level :

1:Atom	2:Atom	3:Atom	4:Atom	5:Atom	6:Atom	7:CLASS	8:CLASS	9:CLASS	12:Atom	13:Atom											
14:Atom	15:Atom	16:Atom	17:CLASS	18:CLASS	19:CLASS	20:CLASS	21:CLASS	22:CLASS													
23:CLASS	24:CLASS	25:CLASS	26:CLASS	29:CLASS	30:CLASS	31:CLASS	32:CLASS	33:Atom													
34:Atom	35:Atom	36:Atom	37:Atom	38:Atom	39:CLASS	40:CLASS	41:CLASS	42:Atom	43:Atom												
44:Atom	45:Atom	46:Atom	47:CLASS	48:CLASS	49:CLASS	50:CLASS	51:CLASS	52:CLASS													
53:CLASS	54:CLASS	55:CLASS	56:CLASS	59:CLASS	60:CLASS	61:CLASS															

fragments assigned product role:

containing 33

fragments assigned reactant/reagent role:

containing 1

=> S 14 SSS SAM

SAMPLE SEARCH INITIATED 18:34:29 FILE 'CASREACT'
SCREENING COMPLETE - 0 REACTIONS TO VERIFY FROM

0 DOCUMENTS

100.0% DONE 0 VERIFIED 0 HIT RXNS 0 DOCS
SEARCH TIME: 00.00.01

FULL FILE PROJECTIONS: ONLINE **COMPLETE**
BATCH **COMPLETE**

PROJECTED VERIFICATIONS: 0 TO 0

PROJECTED ANSWERS: 0 TO 0

L5 0 SEA SSS SAM L4 (0 REACTIONS)

=> S 14 SSS full

FULL SEARCH INITIATED 18:34:37 FILE 'CASREACT'
SCREENING COMPLETE - 0 REACTIONS TO VERIFY FROM

0 DOCUMENTS

100.0% DONE 0 VERIFIED 0 HIT RXNS 0 DOCS
SEARCH TIME: 00.00.01

L6 0 SEA SSS FUL L4 (0 REACTIONS)

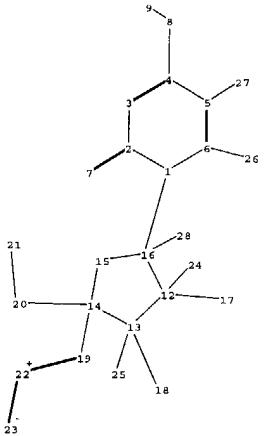
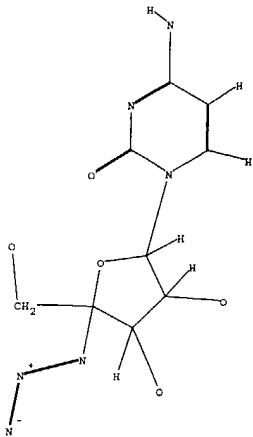
=> d his

(FILE 'HOME' ENTERED AT 18:30:03 ON 10 AUG 2004)

FILE 'REGISTRY' ENTERED AT 18:30:17 ON 10 AUG 2004

FILE 'CASREACT' ENTERED AT 18:30:22 ON 10 AUG 2004

L1 STRUCTURE UPLOADED
L2 0 S L1
L3 0 S L1 SSS FULL
L4 STRUCTURE UPLOADED
L5 0 S L4 SSS SAM
L6 0 S L4 SSS FULL



chain nodes :

7 8 9 17 18 19 20 21 22 23 24 25 26 27 28

ring nodes :

1 2 3 4 5 6 12 13 14 15 16

chain bonds :

1-16 2-7 4-8 5-27 6-26 8-9 12-17 12-24 13-18 13-25 14-19 14-20 16-28 19-22
20-21 22-23

ring bonds :

1-2 1-6 2-3 3-4 4-5 5-6 12-13 12-16 13-14 14-15 15-16

exact/norm bonds :

1-2 1-6 1-16 2-3 2-7 3-4 4-5 4-8 5-6 12-13 12-16 12-17 13-14 13-18 14-15
14-19 15-16 19-22 22-23

exact bonds :

5-27 6-26 8-9 12-24 13-25 14-20 16-28 20-21

G1:O,S,N

G2:H

Match level :

1:Atom 2:Atom 3:Atom 4:Atom 5:Atom 6:Atom 7:CLASS 8:CLASS 9:CLASS 12:Atom 13:Atom
14:Atom 15:Atom 16:Atom 17:CLASS 18:CLASS 19:CLASS 20:CLASS 21:CLASS 22:CLASS
23:CLASS 24:CLASS 25:CLASS 26:CLASS 27:CLASS 28:CLASS